Rechecked SEQ SEARCH results in combination with focus on text search: Bacillus stearothermophilus (STIC)

Item Name								Download Content			
us-10-797-262-1.rge											
us-10-797-262-1.rng											
us-10-797-262-1.rni											
us-10-797-262-1.rnpbm											
us-10-797-262-1.rnpbn											
us-10-797-262-1.rnpm											
us-10-797-262-1.rnpn											
us-10-797-262-1.rst											
us-10-797-262-2.rag											
us-10-797-262-2.rai											
us-10-797-262-2.rapbm											
us-10-797-262-2.rapbn											
us-10-797-262-2.rapm											
us-10-797-262-2.rapn											
us-10-797-262-2.rpr											
us-10-797-262-2.rup											

```
RESULT 1
US-10-797-262-2
; Sequence 2, Application US/10797262
; GENERAL INFORMATION:
; APPLICANT: East Tennessee State University Research Foundation
; APPLICANT: Lampson, Bert C.
; APPLICANT: Velore, Jashree
; TITLE OF INVENTION: RNA-Dependent DNA Polymerase from thermophilic Geobacillus
; TITLE OF INVENTION: stearothermophilus
; FILE REFERENCE: 2826067-000002
; CURRENT APPLICATION NUMBER: US/10/797,262
; CURRENT FILING DATE: 2004-03-10
  NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
   LENGTH: 420
   TYPE: PRT
   ORGANISM: Geobacillus stearothermophilus
US-10-797-262-2
  Query Match
                         100.0%; Score 2173; DB 37; Length 420;
  Best Local Similarity 100.0%; Pred. No. 1.1e-193;
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Matches 420; Conservative 0; Mismatches 0; Indels 0; Gaps

0;

Db	1	MALLERILARDNLITALKRVEANQGAPGIDGVSTDQLRDYIRAHWSTIRAQLLAGTYRPA 60
Qy	61	PVRRVGIPKPGGGTRQLGIPTVVDRLIQQAILQELTPIFDPDFSPSSFGFRPGRNAHDAV 120
Db	61	PVRRVGIPKPGGGTRQLGIPTVVDRLIQQAILQELTPIFDPDFSPSSFGFRPGRNAHDAV 120
Qy	121	${\tt RQAQGYIQEGYRYVVDMDLEKFFDRVNHDILMSRVARKVKDKRVLKLIRAYLQAGVMIEG~180}$
Db	121	
Qy	181	${\tt VKVQTEEGTPQGGPLSPLLANILLDDLDKELEKRGLKFCRYADDCNIYVKSLRAGQRVKQ~240}$
Db	181	VKVQTEEGTPQGGPLSPLLANILLDDLDKELEKRGLKFCRYADDCNIYVKSLRAGQRVKQ 240
Qy	241	SIQRFLEKTLKLKVNEEKSAVDRPWKRAFLGFSFTPERKARIRLAPRSIQRLKQRIRQLT 300
Db	241	
Qy	301	NPNWSISMPERIHRVNQYVMGWIGYFRLVETPSVLQTIEGWIRRRLRLCQWLQWKRVRTR 360
Db	301	
Qy	361	IRELRALGLKETAVMEIANTRKGAWRTTKTPQLHQALGKTYWTAQGLKSLTQRYFELRQG 420
Db	361	IRELRALGLKETAVMEIANTRKGAWRTTKTPQLHQALGKTYWTAQGLKSLTQRYFELRQG 420